



## SEQUENCE LISTING

<110> Sung, Wing  
<120> Xylanases with Enhanced Thermophilicity and Alkalophilicity  
<130> 07121.0003U1  
<140> 09/990,874  
<141> 2001-11-21  
<160> 71  
<170> PatentIn version 3.0  
<210> 1  
<211> 184  
<212> PRT  
<213> Aspergillus niger

<400> 1

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ser | Ala | Gly | Ile | Asn | Tyr | Val | Gln | Asn | Tyr | Asn | Gly | Asn | Leu | Gly | Asp |
| 1   |     |     |     | 5   |     |     |     |     | 10  |     |     |     |     | 15  |     |
| Phe | Thr | Tyr | Asp | Glu | Ser | Ala | Gly | Thr | Phe | Ser | Met | Tyr | Trp | Glu | Asp |
|     |     |     | 20  |     |     |     |     | 25  |     |     |     |     | 30  |     |     |
| Gly | Val | Ser | Ser | Asp | Phe | Val | Val | Gly | Leu | Gly | Trp | Thr | Thr | Gly | Ser |
|     |     | 35  |     |     |     |     | 40  |     |     |     |     | 45  |     |     |     |
| Ser | Asn | Ala | Ile | Thr | Tyr | Ser | Ala | Glu | Tyr | Ser | Ala | Ser | Gly | Ser | Ser |
|     | 50  |     |     |     |     | 55  |     |     |     |     | 60  |     |     |     |     |
| Ser | Tyr | Leu | Ala | Val | Tyr | Gly | Trp | Val | Asn | Tyr | Pro | Gly | Ala | Glu | Tyr |
| 65  |     |     |     | 70  |     |     |     |     | 75  |     |     |     |     | 80  |     |
| Tyr | Ile | Val | Glu | Asp | Tyr | Gly | Asp | Tyr | Asn | Pro | Cys | Ser | Ser | Ala | Thr |
|     |     |     | 85  |     |     |     |     | 90  |     |     |     |     |     | 95  |     |
| Ser | Leu | Gly | Thr | Val | Tyr | Ser | Asp | Gly | Ser | Thr | Tyr | Gln | Val | Cys | Thr |
|     |     |     | 100 |     |     |     |     | 105 |     |     |     |     | 110 |     |     |
| Asp | Thr | Arg | Ile | Asn | Glu | Pro | Ser | Ile | Thr | Gly | Thr | Ser | Thr | Phe | Thr |
|     |     | 115 |     |     |     |     | 120 |     |     |     |     | 125 |     |     |     |
| Gln | Tyr | Phe | Ser | Val | Arg | Glu | Ser | Thr | Arg | Thr | Ser | Gly | Thr | Val | Thr |
|     | 130 |     |     |     |     | 135 |     |     |     |     | 140 |     |     |     |     |
| Val | Ala | Asn | His | Phe | Asn | Phe | Trp | Ala | Gln | His | Gly | Phe | Gly | Asn | Ser |
| 145 |     |     |     | 150 |     |     |     |     |     | 155 |     |     |     | 160 |     |
| Asp | Phe | Asn | Tyr | Gln | Val | Met | Ala | Val | Glu | Ala | Trp | Ser | Gly | Ala | Gly |
|     |     |     |     | 165 |     |     |     |     | 170 |     |     |     |     | 175 |     |

Ser Ala Ser Val Thr Ile Ser Ser  
180

<210> 2

<211> 185

<212> PRT

<213> *Aspergillus tubigensis*

<400> 2

Ser Ala Gly Ile Asn Tyr Val Gln Asn Tyr Asn Gln Asn Leu Gly Asp  
1 5 10 15

Phe Thr Tyr Asp Glu Ser Ala Gly Thr Phe Ser Met Tyr Trp Glu Asp  
20 25 30

Gly Val Ser Ser Asp Phe Val Val Gly Leu Gly Gly Trp Thr Thr Gly  
35 40 45

Ser Ser Asn Ala Ile Thr Tyr Ser Ala Glu Tyr Ser Ala Ser Gly Ser  
50 55 60

Ala Ser Tyr Leu Ala Val Tyr Gly Trp Val Asn Tyr Pro Gln Ala Glu  
65 70 75 80

Tyr Tyr Ile Val Glu Asp Tyr Gly Asp Tyr Asn Pro Cys Ser Ser Ala  
85 90 95

Thr Ser Leu Gly Thr Val Tyr Ser Asp Gly Ser Thr Tyr Gln Val Cys  
100 105 110

Thr Asp Thr Arg Ile Asn Glu Pro Ser Ile Thr Gly Thr Ser Thr Phe  
115 120 125

Thr Gln Tyr Phe Ser Val Arg Glu Ser Thr Arg Thr Ser Gly Thr Val  
130 135 140

Thr Val Ala Asn His Phe Asn Phe Trp Ala His His Gly Phe His Asn  
145 150 155 160

Ser Asp Phe Asn Tyr Gln Val Val Ala Val Glu Ala Trp Ser Gly Ala  
165 170 175

Gly Ser Ala Ala Val Thr Ile Ser Ser  
180 185

<210> 3

<211> 185

<212> PRT

<213> *Bacillus circulans*

<400> 3

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ala | Ser | Thr | Asp | Tyr | Trp | Gln | Asn | Trp | Thr | Asp | Gly | Gly | Gly | Ile | Val | 1   | 5   | 10  | 15  |
| Asn | Ala | Val | Asn | Gly | Ser | Gly | Gly | Asn | Tyr | Ser | Val | Asn | Trp | Ser | Asn | 20  | 25  | 30  |     |
| Thr | Gly | Asn | Phe | Val | Val | Gly | Lys | Gly | Trp | Thr | Thr | Gly | Ser | Pro | Phe | 35  | 40  | 45  |     |
| Arg | Thr | Ile | Asn | Tyr | Asn | Ala | Gly | Val | Trp | Ala | Pro | Asn | Gly | Asn | Gly | 50  | 55  | 60  |     |
| Tyr | Leu | Thr | Leu | Tyr | Gly | Trp | Thr | Arg | Ser | Pro | Leu | Ile | Glu | Tyr | Tyr | 65  | 70  | 75  | 80  |
| Val | Val | Asp | Ser | Trp | Gly | Thr | Tyr | Arg | Pro | Thr | Gly | Thr | Tyr | Lys | Gly | 85  | 90  | 95  |     |
| Thr | Val | Lys | Ser | Asp | Gly | Gly | Thr | Tyr | Asp | Ile | Tyr | Thr | Thr | Thr | Arg | 100 | 105 | 110 |     |
| Tyr | Asn | Ala | Pro | Ser | Ile | Asp | Gly | Asp | Arg | Thr | Thr | Phe | Thr | Gln | Tyr | 115 | 120 | 125 |     |
| Trp | Ser | Val | Arg | Gln | Ser | Lys | Arg | Pro | Thr | Gly | Ser | Asn | Ala | Thr | Ile | 130 | 135 | 140 |     |
| Thr | Phe | Thr | Asn | His | Val | Asn | Ala | Trp | Lys | Ser | His | Gly | Met | Asn | Leu | 145 | 150 | 155 | 160 |
| Gly | Ser | Asn | Trp | Ala | Tyr | Gln | Val | Met | Ala | Thr | Glu | Gly | Tyr | Gln | Ser | 165 | 170 | 175 |     |
| Ser | Gly | Ser | Ser | Asn | Val | Thr | Val | Trp | 180 | 185 |     |     |     |     |     |     |     |     |     |

<210> 4

<211> 201

<212> PRT

<213> Bacillus pumilus

<400> 4

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |    |    |    |    |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|----|----|----|----|
| Arg | Thr | Ile | Thr | Asn | Asn | Glu | Met | Gly | Asn | His | Ser | Gly | Tyr | Asp | Tyr | 1  | 5  | 10 | 15 |
| Glu | Leu | Trp | Lys | Asp | Tyr | Gly | Asn | Thr | Ser | Met | Thr | Leu | Asn | Asn | Gly | 20 | 25 | 30 |    |
| Gly | Ala | Phe | Ser | Ala | Gly | Trp | Asn | Asn | Ile | Gly | Asn | Ala | Leu | Phe | Arg | 35 | 40 | 45 |    |

ATTORNEY DOCKET NO. 07121.0003U1

Lys Gly Lys Lys Phe Asp Ser Thr Arg Thr His His Gln Leu Gly Asn  
50 55 60

Ile Ser Ile Asn Tyr Asn Ala Ser Phe Asn Pro Ser Gly Asn Ser Tyr  
65 70 75 80

Leu Cys Val Tyr Gly Trp Thr Gln Ser Pro Leu Ala Glu Tyr Tyr Ile  
85 90 95

Val Asp Ser Trp Gly Thr Tyr Arg Pro Thr Gly Ala Tyr Lys Gly Ser  
100 105 110

Phe Tyr Ala Asp Gly Gly Thr Tyr Asp Ile Tyr Glu Thr Thr Arg Val  
115 120 125

Asn Gln Pro Ser Ile Ile Gly Ile Ala Thr Phe Lys Gln Tyr Trp Ser  
130 135 140

Val Arg Gln Thr Lys Arg Thr Ser Gly Thr Val Ser Val Ser Ala His  
145 150 155 160

Phe Arg Lys Trp Glu Ser Leu Gly Met Pro Met Gly Lys Met Tyr Glu  
165 170 175

Thr Ala Phe Thr Val Glu Gly Tyr Gln Ser Ser Gly Ser Ala Asn Val  
180 185 190

Met Thr Asn Gln Leu Phe Ile Gly Asn  
195 200

<210> 5

<211> 185

<212> PRT

<213> Bacillus subtilus

<400> 5

Ala Ser Thr Asp Tyr Trp Gln Asn Trp Thr Asp Gly Gly Gly Ile Val  
1 5 10 15

Asn Ala Val Asn Gly Ser Gly Gly Asn Tyr Ser Val Asn Trp Ser Asn  
20 25 30

Thr Gly Asn Phe Val Val Gly Lys Gly Trp Thr Thr Gly Ser Pro Phe  
35 40 45

Arg Thr Ile Asn Tyr Asn Ala Gly Val Trp Ala Pro Asn Gly Asn Gly  
50 55 60

Tyr Leu Thr Leu Tyr Gly Trp Thr Arg Ser Pro Leu Ile Glu Tyr Tyr  
65 70 75 80

Val Val Asp Ser Trp Gly Thr Tyr Arg Pro Thr Gly Thr Tyr Lys Gly  
85 90 95

Thr Val Lys Ser Asp Gly Gly Thr Tyr Asp Ile Tyr Thr Thr Thr Arg  
100 105 110

Tyr Asn Ala Pro Ser Ile Asp Gly Asp Arg Thr Thr Phe Thr Gln Tyr  
115 120 125

Trp Ser Val Arg Gln Ser Lys Arg Pro Thr Gly Ser Asn Ala Thr Ile  
130 135 140

Thr Phe Ser Asn His Val Asn Ala Trp Lys Ser His Gly Met Asn Leu  
145 150 155 160

Gly Ser Asn Trp Ala Tyr Gln Val Met Ala Thr Glu Gly Tyr Gln Ser  
165 170 175

Ser Gly Ser Ser Asn Val Thr Val Trp  
180 185

<210> 6

<211> 211

<212> PRT

<213> Clostridium acetobutylicum

<400> 6

Ser Ala Phe Asn Thr Gln Ala Ala Pro Lys Thr Ile Thr Ser Asn Glu  
1 5 10 15

Ile Gly Val Asn Gly Gly Tyr Asp Tyr Glu Leu Trp Lys Asp Tyr Gly  
20 25 30

Asn Thr Ser Met Thr Leu Lys Asn Gly Gly Ala Phe Ser Cys Gln Trp  
35 40 45

Ser Asn Ile Gly Asn Ala Leu Phe Arg Lys Gly Lys Lys Phe Asn Asp  
50 55 60

Thr Gln Thr Tyr Lys Gln Leu Gly Asn Ile Ser Val Asn Tyr Asn Cys  
65 70 75 80

Asn Tyr Gln Pro Tyr Gly Asn Ser Tyr Leu Cys Val Tyr Gly Trp Thr  
85 90 95

Ser Ser Pro Leu Val Glu Tyr Tyr Ile Val Asp Ser Trp Gly Ser Trp  
100 105 110

Arg Pro Pro Gly Gly Thr Ser Lys Gly Thr Ile Thr Val Asp Gly Gly  
115 120 125

Ile Tyr Asp Ile Tyr Glu Thr Thr Arg Ile Asn Gln Pro Ser Ile Gln  
130 135 140

Gly Asn Thr Thr Phe Lys Gln Tyr Trp Ser Val Arg Arg Thr Lys Arg  
145 150 155 160

Thr Ser Gly Thr Ile Ser Val Ser Lys His Phe Ala Ala Trp Glu Ser  
165 170 175

Lys Gly Met Pro Leu Gly Lys Met His Glu Thr Ala Phe Asn Ile Glu  
180 185 190

Gly Tyr Gln Ser Ser Gly Lys Ala Asp Val Asn Ser Met Ser Ile Asn  
195 200 205

Ile Gly Lys  
210

<210> 7

<211> 206

<212> PRT

<213> Clostridium stercocrarium

<400> 7

Gly Arg Ile Ile Tyr Asp Asn Glu Thr Gly Thr His Gly Gly Tyr Asp  
1 5 10 15

Tyr Glu Leu Trp Lys Asp Tyr Gly Asn Thr Ile Met Glu Leu Asn Asp  
20 25 30

Gly Gly Thr Phe Ser Cys Gln Trp Ser Asn Ile Gly Asn Ala Leu Phe  
35 40 45

Arg Lys Gly Arg Lys Phe Asn Ser Asp Lys Thr Tyr Gln Glu Leu Gly  
50 55 60

Asp Ile Val Val Glu Tyr Gly Cys Asp Tyr Asn Pro Asn Gly Asn Ser  
65 70 75 80

Tyr Leu Cys Val Tyr Gly Trp Thr Arg Asn Phe Leu Val Glu Tyr Tyr  
85 90 95

Ile Val Glu Ser Trp Gly Ser Trp Arg Pro Pro Gly Ala Thr Pro Lys  
100 105 110

Gly Thr Ile Thr Gln Trp Met Ala Gly Thr Tyr Glu Ile Tyr Glu Thr  
115 120 125

Thr Arg Val Asn Gln Pro Ser Ile Asp Gly Thr Ala Thr Phe Gln Gln  
130 135 140

Tyr Trp Ser Val Arg Thr Ser Lys Arg Thr Ser Gly Thr Ile Ser Val  
145 150 155 160

Thr Glu His Phe Lys Gln Trp Glu Arg Met Gly Met Arg Met Gly Lys  
165 170 175

Met Tyr Glu Val Ala Leu Thr Val Glu Gly Tyr Gln Ser Ser Gly Tyr  
180 185 190

Ala Asn Val Tyr Lys Asn Glu Ile Arg Ile Gly Ala Asn Pro  
195 200 205

<210> 8

<211> 211

<212> PRT

<213> Ruminococcus flavefaciens

<400> 8

Ser Ala Ala Asp Gln Gln Thr Arg Gly Asn Val Gly Gly Tyr Asp Tyr  
1 5 10 15

Glu Met Trp Asn Gln Asn Gly Gln Gly Gln Ala Ser Met Asn Pro Gly  
20 25 30

Ala Gly Ser Phe Thr Cys Ser Trp Ser Asn Ile Glu Asn Phe Leu Ala  
35 40 45

Arg Met Gly Lys Asn Tyr Asp Ser Gln Lys Lys Asn Tyr Lys Ala Phe  
50 55 60

Gly Asn Ile Val Leu Thr Tyr Asp Val Glu Tyr Thr Pro Arg Gly Asn  
65 70 75 80

Ser Tyr Met Cys Val Tyr Gly Trp Thr Arg Asn Pro Leu Met Glu Tyr  
85 90 95

Tyr Ile Val Glu Gly Trp Gly Asp Trp Arg Pro Pro Gly Asn Asp Gly  
100 105 110

Glu Val Lys Gly Thr Val Ser Ala Asn Gly Asn Thr Tyr Asp Ile Arg  
115 120 125

Lys Thr Met Arg Tyr Asn Gln Pro Ser Leu Asp Gly Thr Ala Thr Phe  
130 135 140

Pro Gln Tyr Trp Ser Val Arg Gln Thr Ser Gly Ser Ala Asn Asn Gln  
145 150 155 160

Thr Asn Tyr Met Lys Gly Thr Ile Asp Val Ser Lys His Phe Asp Ala  
165 170 175

Trp Ser Ala Ala Gly Leu Asp Met Ser Gly Thr Leu Tyr Glu Val Ser  
180 185 190

Leu Asn Ile Glu Gly Tyr Arg Ser Asn Gly Ser Ala Asn Val Lys Ser  
195 200 205

Val Ser Val  
210

<210> 9

<211> 197

<212> PRT

<213> Schizophyllum cimmune

<400> 9

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ser | Gly | Thr | Pro | Ser | Ser | Thr | Gly | Thr | Asp | Gly | Gly | Tyr | Tyr | Tyr | Ser |
| 1   |     |     |     | 5   |     |     |     |     | 10  |     |     |     |     | 15  |     |
|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
| Trp | Trp | Thr | Asp | Gly | Ala | Gly | Asp | Ala | Thr | Tyr | Gln | Asn | Asn | Gly | Gly |
|     |     |     | 20  |     |     |     |     | 25  |     |     |     |     | 30  |     |     |
|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
| Gly | Ser | Tyr | Thr | Leu | Thr | Trp | Ser | Gly | Asn | Asn | Gly | Asn | Leu | Val | Gly |
|     | 35  |     |     |     |     |     | 40  |     |     |     |     | 45  |     |     |     |
|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
| Gly | Lys | Gly | Trp | Asn | Pro | Gly | Ala | Ala | Ser | Arg | Ser | Ile | Ser | Tyr | Ser |
|     | 50  |     |     |     |     | 55  |     |     |     |     | 60  |     |     |     |     |
|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
| Gly | Thr | Tyr | Gln | Pro | Asn | Gly | Asn | Ser | Tyr | Leu | Ser | Val | Tyr | Gly | Trp |
| 65  |     |     |     |     | 70  |     |     |     |     | 75  |     |     |     |     | 80  |
|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
| Thr | Arg | Ser | Ser | Leu | Ile | Glu | Tyr | Tyr | Ile | Val | Glu | Ser | Tyr | Gly | Ser |
|     |     |     |     | 85  |     |     |     |     | 90  |     |     |     |     | 95  |     |
|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
| Tyr | Asp | Pro | Ser | Ser | Ala | Ala | Ser | His | Lys | Gly | Ser | Val | Thr | Cys | Asn |
|     |     |     | 100 |     |     |     |     | 105 |     |     |     |     |     | 110 |     |
|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
| Gly | Ala | Thr | Tyr | Asp | Ile | Leu | Ser | Thr | Trp | Arg | Tyr | Asn | Ala | Pro | Ser |
|     |     | 115 |     |     |     |     |     | 120 |     |     |     |     | 125 |     |     |
|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
| Ile | Asp | Gly | Thr | Gln | Thr | Phe | Glu | Gln | Phe | Trp | Ser | Val | Arg | Asn | Pro |
|     | 130 |     |     |     |     | 135 |     |     |     |     | 140 |     |     |     |     |
|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
| Lys | Lys | Ala | Pro | Gly | Gly | Ser | Ile | Ser | Gly | Thr | Val | Asp | Val | Gln | Cys |
| 145 |     |     |     |     | 150 |     |     |     |     | 155 |     |     |     |     | 160 |
|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
| His | Phe | Asp | Ala | Trp | Lys | Gly | Leu | Gly | Met | Asn | Leu | Gly | Ser | Glu | His |
|     |     |     |     | 165 |     |     |     |     | 170 |     |     |     |     | 175 |     |
|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
| Asn | Tyr | Gln | Ile | Val | Ala | Thr | Glu | Gly | Tyr | Gln | Ser | Ser | Gly | Thr | Ala |
|     |     |     | 180 |     |     |     |     | 185 |     |     |     |     |     | 190 |     |
|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
| Thr | Ile | Thr | Val | Thr |     |     |     |     |     |     |     |     |     |     |     |
|     |     |     |     | 195 |     |     |     |     |     |     |     |     |     |     |     |

<210> 10

<211> 191

<212> PRT

<213> Streptomyces lividans Xyl B



<400> 10

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Asp | Thr | Val | Val | Thr | Thr | Asn | Gln | Glu | Gly | Thr | Asn | Asn | Gly | Tyr | Tyr | 1   | 5   | 10  | 15  |
| Tyr | Ser | Phe | Trp | Thr | Asp | Ser | Gln | Gly | Thr | Val | Ser | Met | Asn | Met | Gly | 20  | 25  | 30  |     |
| Ser | Gly | Gly | Gln | Tyr | Ser | Thr | Ser | Trp | Arg | Asn | Thr | Gly | Asn | Phe | Val | 35  | 40  | 45  |     |
| Ala | Gly | Lys | Gly | Trp | Ala | Asn | Gly | Gly | Arg | Arg | Thr | Val | Gln | Tyr | Ser | 50  | 55  | 60  |     |
| Gly | Ser | Phe | Asn | Pro | Ser | Gly | Asn | Ala | Tyr | Leu | Ala | Leu | Tyr | Gly | Trp | 65  | 70  | 75  | 80  |
| Thr | Ser | Asn | Pro | Leu | Val | Glu | Tyr | Tyr | Ile | Val | Asp | Asn | Trp | Gly | Thr | 85  | 90  | 95  |     |
| Tyr | Arg | Pro | Thr | Gly | Glu | Tyr | Lys | Gly | Thr | Val | Thr | Ser | Asp | Gly | Gly | 100 | 105 | 110 |     |
| Thr | Tyr | Asp | Ile | Tyr | Lys | Thr | Thr | Arg | Val | Asn | Lys | Pro | Ser | Val | Glu | 115 | 120 | 125 |     |
| Gly | Thr | Arg | Thr | Phe | Asp | Gln | Tyr | Trp | Ser | Val | Arg | Gln | Ser | Lys | Arg | 130 | 135 | 140 |     |
| Thr | Gly | Gly | Thr | Ile | Thr | Thr | Gly | Asn | His | Phe | Asp | Ala | Trp | Ala | Arg | 145 | 150 | 155 | 160 |
| Ala | Gly | Met | Pro | Leu | Gly | Asn | Phe | Ser | Tyr | Tyr | Met | Ile | Asn | Ala | Thr | 165 | 170 | 175 |     |
| Glu | Gly | Tyr | Gln | Ser | Ser | Gly | Thr | Ser | Ser | Ile | Asn | Val | Gly | Gly | 180 | 185 | 190 |     |     |

<210> 11

<211> 191

<212> PRT

<213> Streptomyces lividans Xyl C

<400> 11

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |    |    |    |    |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|----|----|----|----|
| Ala | Thr | Thr | Ile | Thr | Thr | Asn | Gln | Thr | Gly | Thr | Asp | Gly | Met | Tyr | Tyr | 1  | 5  | 10 | 15 |
| Ser | Phe | Trp | Thr | Asp | Gly | Gly | Gly | Ser | Val | Ser | Met | Thr | Leu | Asn | Gly | 20 | 25 | 30 |    |
| Gly | Gly | Ser | Tyr | Ser | Thr | Gln | Trp | Thr | Asn | Cys | Gly | Asn | Phe | Val | Ala | 35 | 40 | 45 |    |

Gly Lys Gly Trp Ser Thr Gly Asp Gly Asn Val Arg Tyr Asn Gly Tyr  
50 55 60

Phe Asn Pro Val Gly Asn Gly Tyr Gly Cys Leu Tyr Gly Trp Thr Ser  
65 70 75 80

Asn Pro Leu Val Glu Tyr Tyr Ile Val Asp Asn Trp Gly Ser Tyr Arg  
85 90 95

Pro Thr Gly Thr Tyr Lys Gly Thr Val Ser Ser Asp Gly Gly Thr Tyr  
100 105 110

Asp Ile Tyr Gln Thr Thr Arg Tyr Asn Ala Pro Ser Val Glu Gly Thr  
115 120 125

Lys Thr Phe Gln Gln Tyr Trp Ser Val Arg Gln Ser Lys Val Thr Ser  
130 135 140

Gly Ser Gly Thr Ile Thr Thr Gly Asn His Phe Asp Ala Trp Ala Arg  
145 150 155 160

Ala Gly Met Asn Met Gly Gln Phe Arg Tyr Tyr Met Ile Asn Ala Thr  
165 170 175

Glu Gly Tyr Gln Ser Ser Gly Ser Ser Asn Ile Thr Val Ser Gly  
180 185 190

<210> 12

<211> 189

<212> PRT

<213> Streptomyces sp. No. 36a

<400> 12

Ala Thr Thr Ile Thr Asn Glu Thr Gly Tyr Asp Gly Met Tyr Tyr Ser  
1 5 10 15

Phe Trp Thr Asp Gly Gly Gly Ser Val Ser Met Thr Leu Asn Gly Gly  
20 25 30

Gly Ser Tyr Ser Thr Arg Trp Thr Asn Cys Gly Asn Phe Val Ala Gly  
35 40 45

Lys Gly Trp Ala Asn Gly Gly Arg Arg Thr Val Arg Tyr Thr Gly Trp  
50 55 60

Phe Asn Pro Ser Gly Asn Gly Tyr Gly Cys Leu Tyr Gly Trp Thr Ser  
65 70 75 80

Asn Pro Leu Val Glu Tyr Tyr Ile Val Asp Asn Trp Gly Ser Tyr Arg  
85 90 95

Pro Thr Gly Glu Thr Arg Gly Thr Val His Ser Asp Gly Gly Thr Tyr  
100 105 110

Asp Ile Tyr Lys Thr Thr Arg Tyr Asn Ala Pro Ser Val Glu Ala Pro  
115 120 125

Ala Ala Phe Asp Gln Tyr Trp Ser Val Arg Gln Ser Lys Val Thr Ser  
130 135 140

Gly Thr Ile Thr Thr Gly Asn His Phe Asp Ala Trp Ala Arg Ala Gly  
145 150 155 160

Met Asn Met Gly Asn Phe Arg Tyr Tyr Met Ile Asn Ala Thr Glu Gly  
165 170 175

Tyr Gln Ser Ser Gly Ser Ser Thr Ile Thr Val Ser Gly  
180 185

<210> 13

<211> 189

<212> PRT

<213> Thermomonospora fusca

<400> 13

Ala Val Thr Ser Asn Glu Thr Gly Tyr His Asp Gly Tyr Phe Tyr Ser  
1 5 10 15

Phe Trp Thr Asp Ala Pro Gly Thr Val Ser Met Glu Leu Gly Pro Gly  
20 25 30

Gly Asn Tyr Ser Thr Ser Trp Arg Asn Thr Gly Asn Phe Val Ala Gly  
35 40 45

Lys Gly Trp Ala Thr Gly Gly Arg Arg Thr Val Thr Tyr Ser Ala Ser  
50 55 60

Phe Asn Pro Ser Gly Asn Ala Tyr Leu Thr Leu Tyr Gly Trp Thr Arg  
65 70 75 80

Asn Pro Leu Val Glu Tyr Tyr Ile Val Glu Ser Trp Gly Thr Tyr Arg  
85 90 95

Pro Thr Gly Thr Tyr Met Gly Thr Val Thr Thr Asp Gly Gly Thr Tyr  
100 105 110

Asp Ile Tyr Lys Thr Thr Arg Tyr Asn Ala Pro Ser Ile Glu Gly Thr  
115 120 125

Arg Thr Phe Asp Gln Tyr Trp Ser Val Arg Gln Ser Lys Arg Thr Ser  
130 135 140

Gly Thr Ile Thr Ala Gly Asn His Phe Asp Ala Trp Ala Arg His Gly  
145 150 155 160

Met His Leu Gly Thr His Asp Tyr Met Ile Met Ala Thr Glu Gly Tyr  
165 170 175

Gln Ser Ser Gly Ser Ser Asn Val Thr Leu Gly Thr Ser  
180 185

<210> 14

<211> 190

<212> PRT

<213> Trichoderma harzanium

<400> 14

Gln Thr Ile Gly Pro Gly Thr Gly Tyr Ser Asn Gly Tyr Tyr Tyr Ser  
1 5 10 15  
Tyr Trp Asn Asp Gly His Ala Gly Val Thr Tyr Thr Asn Gly Gly Gly  
20 25 30  
Gly Ser Phe Thr Val Asn Trp Ser Asn Ser Gly Asn Phe Val Gly Gly  
35 40 45  
Lys Gly Trp Gln Pro Gly Thr Lys Asn Lys Val Ile Asn Phe Ser Gly  
50 55 60  
Ser Tyr Asn Pro Asn Gly Asn Ser Tyr Leu Ser Ile Tyr Gly Trp Ser  
65 70 75 80  
Arg Asn Pro Leu Ile Glu Tyr Tyr Ile Val Glu Asn Phe Gly Thr Tyr  
85 90 95  
Asn Pro Ser Thr Gly Ala Thr Lys Leu Gly Glu Val Thr Ser Asp Gly  
100 105 110  
Ser Val Tyr Asp Ile Tyr Arg Thr Gln Arg Val Asn Gln Pro Ser Ile  
115 120 125  
Ile Gly Thr Ala Thr Phe Tyr Gln Tyr Trp Ser Val Arg Arg Asn His  
130 135 140  
Arg Ser Ser Gly Ser Val Asn Thr Ala Asn His Phe Asn Ala Trp Ala  
145 150 155 160  
Ser His Gly Leu Thr Leu Gly Thr Met Asp Tyr Gln Ile Val Ala Val  
165 170 175  
Glu Gly Tyr Phe Ser Ser Gly Ser Ala Ser Ile Thr Val Ser  
180 185 190

<210> 15

<211> 178

<212> PRT

<213> Trichoderma ressei Xyl I

Ser Asn

<210> 16

<211> 190

<212> PRT

<213> Trichoderma ressei Xyl II

<400> 16

13

Lys Gly Trp Gln Pro Gly Thr Lys Asn Lys Val Ile Asn Phe Ser Gly  
50 55 60

Ser Tyr Asn Pro Asn Gly Asn Ser Tyr Leu Ser Val Tyr Gly Trp Ser  
65 70 75 80

Arg Asn Pro Leu Ile Glu Tyr Tyr Ile Val Glu Asn Phe Gly Thr Tyr  
85 90 95

Asn Pro Ser Thr Gly Ala Thr Lys Leu Gly Glu Val Thr Ser Asp Gly  
100 105 110

Ser Val Tyr Asp Ile Tyr Arg Thr Gln Arg Val Asn Gln Pro Ser Ile  
115 120 125

Ile Gly Thr Ala Thr Phe Tyr Gln Tyr Trp Ser Val Arg Arg Asn His  
130 135 140

Arg Ser Ser Gly Ser Val Asn Thr Ala Asn His Phe Asn Ala Trp Ala  
145 150 155 160

Gln Gln Gly Leu Thr Leu Gly Thr Met Asp Tyr Gln Ile Val Ala Val  
165 170 175

Glu Gly Tyr Phe Ser Ser Gly Ser Ala Ser Ile Thr Val Ser  
180 185 190

<210> 17

<211> 190

<212> PRT

<213> Trichoderma viride

<400> 17

Gln Thr Ile Gln Pro Gly Thr Gly Phe Asn Asn Gly Tyr Phe Tyr Ser  
1 5 10 15

Tyr Trp Asn Asp Gly His Gly Gly Val Thr Tyr Thr Asn Gly Pro Gly  
20 25 30

Gly Gln Phe Ser Val Asn Trp Ser Asn Ser Gly Asn Phe Val Gly Gly  
35 40 45

Lys Gly Trp Gln Pro Gly Thr Lys Asn Lys Val Ile Asn Phe Ser Gly  
50 55 60

Ser Tyr Asn Pro Asn Gly Asn Ser Tyr Leu Ser Val Tyr Gly Trp Ser  
65 70 75 80

Arg Asn Pro Leu Ile Glu Tyr Tyr Ile Val Glu Asn Phe Gly Thr Tyr  
85 90 95

Asn Pro Ser Thr Gly Ala Thr Lys Leu Gly Glu Val Thr Ser Asp Gly  
100 105 110

Ser Val Tyr Asp Ile Tyr Arg Thr Gln Arg Val Asn Gln Pro Ser Ile  
115 120 125

Ile Gly Thr Ala Thr Phe Tyr Gln Tyr Trp Ser Val Arg Arg Thr His  
130 135 140

Arg Ser Ser Gly Ser Val Asn Thr Ala Asn His Phe Asn Ala Trp Ala  
145 150 155 160

Gln Gln Gly Leu Thr Leu Gly Thr Met Asp Tyr Gln Ile Val Ala Val  
165 170 175

Glu Gly Tyr Phe Ser Ser Gly Ser Ala Ser Ile Thr Val Ser  
180 185 190

<210> 18

<211> 202

<212> PRT

<213> Fibrobacter succinognees

<400> 18

Asn Ser Ser Val Thr Gly Asn Val Gly Ser Ser Pro Tyr His Tyr Glu  
1 5 10 15

Ile Trp Tyr Gln Gly Gly Asn Asn Ser Met Thr Phe Tyr Asp Asn Gly  
20 25 30

Thr Tyr Lys Ala Ser Trp Asn Gly Thr Asn Asp Phe Leu Ala Arg Val  
35 40 45

Gly Phe Lys Tyr Asp Glu Lys His Thr Tyr Glu Glu Leu Gly Pro Ile  
50 55 60

Asp Ala Tyr Tyr Lys Trp Ser Lys Gln Gly Ser Ala Gly Gly Tyr Asn  
65 70 75 80

Tyr Ile Gly Ile Tyr Gly Trp Thr Val Asp Pro Leu Val Glu Tyr Tyr  
85 90 95

Ile Val Asp Asp Trp Phe Asn Lys Pro Gly Ala Asn Leu Leu Gly Gln  
100 105 110

Arg Lys Gly Glu Phe Thr Val Asp Gly Asp Thr Tyr Glu Ile Trp Gln  
115 120 125

Asn Thr Arg Val Gln Gln Pro Ser Ile Lys Gly Thr Gln Thr Phe Pro  
130 135 140

Gln Tyr Phe Ser Val Arg Lys Ser Ala Arg Ser Cys Gly His Ile Asp  
145 150 155 160

Ile Thr Ala His Met Lys Lys Trp Glu Glu Leu Gly Met Lys Met Gly  
165 170 175

Lys Met Tyr Glu Ala Lys Val Leu Val Glu Ala Gly Gly Gly Ser Gly  
180 185 190

Ser Phe Asp Val Thr Tyr Phe Lys Met Thr  
195 200

<210> 19

<211> 189

<212> PRT

<213> Asparigillus awamori var. kawachi

<400> 19

Arg Ser Thr Pro Ser Ser Thr Gly Glu Asn Asn Gly Tyr Tyr Tyr Ser  
1 5 10 15

Phe Trp Thr Asp Gly Gly Gly Asp Val Thr Tyr Thr Asn Gly Asn Ala  
20 25 30

Gly Ser Tyr Ser Val Glu Trp Ser Asn Val Gly Asn Phe Val Gly Gly  
35 40 45

Lys Gly Trp Asn Pro Gly Ser Ala Lys Asp Ile Thr Tyr Ser Gly Asn  
50 55 60

Phe Thr Pro Ser Gly Asn Gly Tyr Leu Ser Val Tyr Gly Trp Thr Thr  
65 70 75 80

Asp Pro Leu Ile Glu Tyr Tyr Ile Val Glu Ser Tyr Gly Asp Tyr Asn  
85 90 95

Pro Gly Ser Gly Gly Thr Thr Arg Gly Asn Val Ser Ser Asp Gly Ser  
100 105 110

Val Tyr Asp Ile Tyr Thr Ala Thr Arg Thr Asn Ala Pro Ser Ile Asp  
115 120 125

Gly Thr Gln Thr Phe Ser Gln Tyr Trp Ser Val Arg Gln Asn Lys Arg  
130 135 140

Val Gly Gly Thr Val Thr Thr Ser Asn His Phe Asn Ala Trp Ala Lys  
145 150 155 160

Leu Gly Met Asn Leu Gly Thr His Asn Tyr Gln Ile Leu Ala Thr Glu  
165 170 175

Gly Tyr Gln Ser Ser Gly Ser Ser Ser Ile Thr Ile Gln  
180 185

<210> 20



<211> 194

<212> PRT

<213> Thermomyces lanuginosus

<400> 20

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Gln | Thr | Thr | Pro | Asn | Ser | Glu | Gly | Trp | His | Asp | Gly | Tyr | Tyr | Tyr | Ser | 1   | 5   | 10  | 15  |
| Trp | Trp | Ser | Asp | Gly | Gly | Ala | Gln | Ala | Thr | Tyr | Thr | Asn | Leu | Glu | Gly | 20  | 25  | 30  |     |
| Gly | Thr | Tyr | Glu | Ile | Ser | Trp | Gly | Asp | Gly | Gly | Asn | Leu | Val | Gly | Gly | 35  | 40  | 45  |     |
| Lys | Gly | Trp | Asn | Pro | Gly | Leu | Asn | Ala | Arg | Ala | Ile | His | Phe | Glu | Gly | 50  | 55  | 60  |     |
| Val | Tyr | Gln | Pro | Asn | Gly | Asn | Ser | Tyr | Leu | Ala | Val | Tyr | Gly | Trp | Thr | 65  | 70  | 75  | 80  |
| Arg | Asn | Pro | Leu | Val | Glu | Tyr | Tyr | Ile | Val | Glu | Asn | Phe | Gly | Thr | Tyr | 85  | 90  | 95  |     |
| Asp | Pro | Ser | Ser | Gly | Ala | Thr | Asp | Leu | Gly | Thr | Val | Glu | Cys | Asp | Gly | 100 | 105 | 110 |     |
| Ser | Ile | Tyr | Arg | Leu | Gly | Lys | Thr | Thr | Arg | Val | Asn | Ala | Pro | Ser | Ile | 115 | 120 | 125 |     |
| Asp | Gly | Thr | Gln | Thr | Phe | Asp | Gln | Tyr | Trp | Ser | Val | Arg | Gln | Asp | Lys | 130 | 135 | 140 |     |
| Arg | Thr | Ser | Gly | Thr | Val | Gln | Thr | Gly | Cys | His | Phe | Asp | Ala | Trp | Ala | 145 | 150 | 155 | 160 |
| Arg | Ala | Gly | Leu | Asn | Val | Asn | Gly | Asp | His | Tyr | Tyr | Gln | Ile | Val | Ala | 165 | 170 | 175 |     |
| Thr | Glu | Gly | Tyr | Phe | Ser | Ser | Gly | Tyr | Ala | Arg | Ile | Thr | Val | Ala | Asp | 180 | 185 | 190 |     |
| Val | Gly |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |

<210> 21

<211> 76

<212> DNA

<213> Artificial Sequence

<220>

<223> Trx-1

<400> 21  
ctagctaagg aggctgcaga tgcaaacaat acaaccagga accggttaca acaacggtta 60  
cttttacagc tattgg 76

<210> 22

<211> 78

<212> DNA

<213> Artificial Sequence

<220>

<223> XyTv-2

<400> 22  
aacgatggcc atggtggtgt tacctataca aacggggcccg gaggcccaatt tagcgtcaat 60  
tggtctaact ccggaaac 78

<210> 23

<211> 78

<212> DNA

<213> Artificial Sequence

<220>

<223> Trx-3

<400> 23  
ttcgtaggtg gaaaagggtg gcaacccggg accaaaaata aggtgatcaa cttctctgga 60  
tcttataatc cgaatggg 78

<210> 24

<211> 74

<212> DNA

<213> Artificial Sequence

<220>

<223> XyTv-4

<400> 24  
aattcatact taagcgtcta tggctggtct agaaacccac tgattgaata ttacattgtc 60  
gaaaatttcg gtac 74

<210> 25

<211> 51

<212> DNA

<213> Artificial Sequence

<220>

<223> Trx-8

<400> 25  
gattcctccg acgtctacgt ttgttatggt ggtccttggc caatgttggt g 51

<210> 26

<211> 84

<212> DNA

<213> Artificial Sequence

<220>

<223> XyTv-7

<400> 26  
ccaatgaaaa tgtcgataac cttgctaccg gtaccaccac aatggatatg tttgcccggg 60

cctccgggta aatcgcagtt aacc 84

<210> 27

<211> 78

<212> DNA

<213> Artificial Sequence

<220>

<223> Trx-6

<400> 27  
agattgaggg ctttgaagca tccacctttt ccaaccgttg ggccctgggt tttattccac 60  
tagttgaaga gacctaga 78

<210> 28

<211> 85

<212> DNA

<213> Artificial Sequence

<220>

<223> XyTv-5

<400> 28  
atattagggt tacccttaag tatgaattcg cagataccga ccagatcttt gggtgactaa 60  
cttataatgt aacagctttt aaagc 85

<210> 29

<211> 58

<212> DNA

<213> Artificial Sequence

<220>

<223> XyTv-101

<400> 29  
tcgacaattt cggtacctac aatccgagta ccggcgccac aaaattaggc gaagtcac 58

<210> 30

<211> 53

<212> DNA

<213> Artificial Sequence

<220>

<223> XyTv-102

<400> 30  
tagtgatgga tccgtatatg atatctaccg tacccaacgc gttaatcagc cat 53

<210> 31

<211> 59

<212> DNA

<213> Artificial Sequence

<220>

<223> Trx-103

<400> 31  
cgatcattgg aaccgccacc ttttatcagt actggagtgt tagacgtaat catcggagc 59

<210> 32

<211> 69

<212> DNA

<213> Artificial Sequence

<220>

<223> XyTv-104

<400> 32  
tccggttcgg ttaatactgc gaatcacttt aatgcatggg cacagcaagg gttaacccta 60  
ggtacaatg 69

<210> 33

<211> 67

<212> DNA

<213> Artificial Sequence

<220>

<223> XyTv-105

<400> 33  
gattatcaaa tcgtagcggg ggaaggctac ttctcgagtg gttccgctag tattacagtg 60  
agctaaa 67

<210> 34

<211> 73

<212> DNA

<213> Artificial Sequence

<220>

<223> XyTv-110

<400> 34  
gttaaagcca tggatggttag gctcatggcc gcggtgtttt aatccgcttc agtgatcact 60  
acctaggcat ata 73

<210> 35

<211> 54

<212> DNA

<213> Artificial Sequence

<220>

<223> XyTv-109

<400> 35  
ctatagatgg catgggttgc gcaattagtc ggtagctagt aaccttggcg gtgg 54

<210> 36

<211> 60

<212> DNA

<213> Artificial Sequence

<220>

<223> XyTv-108

<400> 36  
aaaatagtca tgacctcaca atctgcatta gtagcctcga ggccaagcca attatgacgc 60

<210> 37

<211> 66

<212> DNA

<213> Artificial Sequence

<220>

<223> XyTv-107

<400> 37  
ttagtgaaat tacgtacccg tgctggtccc aattgggatc catgttacct aatagtttag 60  
catcgc 66

<210> 38

<211> 53

<212> DNA

<213> Artificial Sequence

<220>

<223> XyTv-106

<400> 38  
caccttccga tgaagagctc accaaggcga tcataatgtc actcgatttc tag 53

<210> 39

<211> 596

<212> DNA

<213> Artificial Sequence

<220>

<223> TrX

<400> 39  
ctagctaagg aggctgcaga tgcaacaat acaaccagga accggttaca acaacggtta 60  
cttttacagc tattggaacg atggccatgg tgggtgtacc tatacaaacg ggcccggagg 120  
ccaatttagc gtcaattggt ctaactccg aaacttcgta ggtggaaaag gttggcaacc 180  
cgggaccaaa aataagggtga tcaacttctc tggatcttat aatccgaatg ggaattcata 240  
cttaagcgtc tatggctggt ctagaaaccc actgattgaa tattacattg tcgaaaattt 300  
cgggtacctac aatccgagta ccggcgccac aaaattaggc gaagtcacta gtgatggatc 360  
cgtatatgat atctaccgta cccaacgagt taatcagcca tcgatcattg gaaccgccac 420  
cttttatcag tactggagtg ttagacgtaa tcatcgagc tccgggtcgg ttaatactgc 480

gaatcacttt aatgcatggg cacagcaagg gttaacccta ggtacaatgg attatcaaat 540  
cgtagcgggtg gaaggctact tctcgagtgg ttccgctagt attacagtga gctaaa 596

<210> 40

<211> 36

<212> DNA

<213> Artificial Sequence

<220>

<223> Tx-75A-1

<400> 40  
tgggaattca tacttagccg tctatggctg gtctag 36

<210> 41

<211> 42

<212> DNA

<213> Artificial Sequence

<220>

<223> Tx-105H-1

<400> 41  
accggcgcca caaaacacgg cgaagtcact agtgatggat cc 42

<210> 42

<211> 44

<212> DNA

<213> Artificial Sequence

<220>

<223> Tx-C1

<400> 42  
ccaaggcgat cataatgtca ctcgatttct agaacttcga accc 44

<210> 43



<211> 36

<212> DNA

<213> Artificial Sequence

<220>

<223> Tx-del(123-144)-1r

<400> 43

cggagctccg acgcgttggg tacggtagat atcata

36

<210> 44

<211> 42

<212> DNA

<213> Artificial Sequence

<220>

<223> Tx-105R-1

<400> 44

accggcgcca caaaaagagg cgaagtcact agtgatggat cc

42

<210> 45

<211> 41

<212> DNA

<213> Artificial Sequence

<220>

<223> Tx-N1

<400> 45

ctagctaagg aggctgcaga tgcaaacaat acaaccagga a

41

<210> 46

<211> 36

<212> DNA

<213> Artificial Sequence

<220>

<223> Tx-75-G1

<400> 46

tgggaattca tacttaggcg tctatggctg gtctag

36

<210> 47

<211> 66

<212> DNA

<213> Artificial Sequence

<220>

<223> Tx-144R-1r

<400> 47

ccatgcatta aagtgattcg cagtattaac cgaaccggag ctccgacgat tacgtctaac

60

actcca

66

<210> 48

<211> 44

<212> DNA

<213> Artificial Sequence

<220>

<223> Tx-161R-1r

<400> 48

gtacctaggg ttaacccttg ccgtgcccac gcattaaagt gatt

44

<210> 49

<211> 40

<212> DNA

<213> Artificial Sequence

<220>

<223> Tx-125A 129E-1

<400> 49  
ccaacgcgtt aatgcgccat cgatcgaggg aaccgccacc 40

<210> 50

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> Tx-116G-1

<400> 50  
gacggatccg tatatggtat ctaccg 26

<210> 51

<211> 36

<212> DNA

<213> Artificial Sequence

<220>

<223> Tx-118C-1

<400> 51  
gacggatccg tatatgatat ctgccgtacc caacgc 36

<210> 52

<211> 39

<212> DNA

<213> Artificial Sequence

<220>

<223> Tx-10H11D-1

<400> 52  
ggaaccgggt accacgacgg ttactttttac agctattgg 39

<210> 53

<211> 36

<212> DNA

<213> Artificial Sequence

<220>

<223> Tx-116G118C-1

<400> 53  
gacggatccg tatatggtat ctgccgtacc caacgc

36

<210> 54

<211> 184

<212> PRT

<213> Aspergillus kawachii

<400> 54

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Ser | Ala | Gly | Ile | Asn | Tyr | Val | Gln | Asn | Tyr | Asn | Gly | Asn | Leu | Ala | Asp | 1   | 5   | 10  | 15  |
| Phe | Thr | Tyr | Asp | Glu | Ser | Ala | Gly | Thr | Phe | Ser | Met | Tyr | Trp | Glu | Asp | 20  | 25  | 30  |     |
| Gly | Val | Ser | Ser | Asp | Phe | Val | Val | Gly | Leu | Gly | Trp | Thr | Thr | Gly | Ser | 35  | 40  | 45  |     |
| Ser | Asn | Ala | Ile | Ser | Tyr | Ser | Ala | Glu | Tyr | Ser | Ala | Ser | Gly | Ser | Ser | 50  | 55  | 60  |     |
| Ser | Tyr | Leu | Ala | Val | Tyr | Gly | Trp | Val | Asn | Tyr | Pro | Gln | Ala | Glu | Tyr | 65  | 70  | 75  | 80  |
| Tyr | Ile | Val | Glu | Asp | Tyr | Gly | Asp | Tyr | Asn | Pro | Cys | Ser | Ser | Ala | Thr | 85  | 90  | 95  |     |
| Ser | Leu | Gly | Thr | Val | Tyr | Ser | Asp | Gly | Ser | Thr | Tyr | Gln | Val | Cys | Thr | 100 | 105 | 110 |     |
| Asp | Thr | Arg | Thr | Asn | Glu | Pro | Ser | Ile | Thr | Gly | Thr | Ser | Thr | Phe | Thr | 115 | 120 | 125 |     |
| Gln | Tyr | Phe | Ser | Val | Arg | Glu | Ser | Thr | Arg | Thr | Ser | Gly | Thr | Val | Thr | 130 | 135 | 140 |     |
| Val | Ala | Asn | His | Phe | Asn | Phe | Trp | Ala | Gln | His | Gly | Phe | Gly | Asn | Ser | 145 | 150 | 155 | 160 |
| Asp | Phe | Asn | Tyr | Gln | Val | Met | Ala | Val | Glu | Ala | Trp | Ser | Gly | Ala | Gly | 165 | 170 | 175 |     |

Ser Ala Ser Val Thr Ile Ser Ser  
180

<210> 55

<211> 190

<212> PRT

<213> Artificial Sequence

<220>

<223> TrX-H-11D-ML-75A105H-118C-125A129E-144R161R (TrX-H-11D-ML-AHCAE-RR)

<400> 55

Gln Thr Ile Gln Pro Gly Thr Gly Tyr His Asp Gly Tyr Phe Tyr Ser  
1 5 10 15

Tyr Trp Asn Asp Gly His Gly Gly Val Thr Met Thr Leu Gly Pro Gly  
20 25 30

Gly Gln Phe Ser Val Asn Trp Ser Asn Ser Gly Asn Phe Val Gly Gly  
35 40 45

Lys Gly Trp Gln Pro Gly Thr Lys Asn Lys Val Ile Asn Phe Ser Gly  
50 55 60

Ser Tyr Asn Pro Asn Gly Asn Ser Tyr Leu Ala Val Tyr Gly Trp Ser  
65 70 75 80

Arg Asn Pro Leu Ile Glu Tyr Tyr Ile Val Glu Asn Phe Gly Thr Tyr  
85 90 95

Asn Pro Ser Thr Gly Ala Thr Lys His Gly Glu Val Thr Ser Asp Gly  
100 105 110

Ser Val Tyr Asp Ile Cys Arg Thr Gln Arg Val Asn Ala Pro Ser Ile  
115 120 125

Glu Gly Thr Ala Thr Phe Tyr Gln Tyr Trp Ser Val Arg Arg Asn Arg  
130 135 140

Arg Ser Ser Gly Ser Val Asn Thr Ala Asn His Phe Asn Ala Trp Ala  
145 150 155 160

Arg Gln Gly Leu Thr Leu Gly Thr Met Asp Tyr Gln Ile Val Ala Val  
165 170 175

Glu Gly Tyr Phe Ser Ser Gly Ser Ala Ser Ile Thr Val Ser  
180 185 190

<210> 56

<211> 112

<212> DNA

<213> Artificial Sequence

<220>

<223> TrX-HML

<400> 56  
ctagctaagg aggctgcaga tgcaaacaat acaaccagga accggttacc acaacggtta 60  
cttttacagc tattggaacg atggccatgg aggcgtcaca atgactctgg gg 112

<210> 57

<211> 30

<212> PRT

<213> Artificial Sequence

<220>

<223> TrX-HML

<400> 57

Gln Thr Ile Gln Pro Gly Thr Gly Tyr His Asn Gly Tyr Phe Tyr Ser  
1 5 10 15

Tyr Trp Asn Asp Gly His Gly Gly Val Thr Met Thr Leu Gly  
20 25 30

<210> 58

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> TX-105R-1

<400> 58

Thr Gly Ala Thr Lys Arg Gly Glu Val Thr Ser Asp Gly Ser  
1 5 10

<210> 59

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> TX-C1

<400> 59

Gly Ser Ala Ser Ile Thr Val Ser  
1 5

<210> 60

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<223> TX-75A-1

<400> 60

Asn Gly Asn Ser Tyr Leu Ala Val Tyr Gly Trp Ser Arg  
1 5 10

<210> 61

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<223> TX-75G-1

<400> 61

Asn Gly Asn Ser Tyr Leu Gly Val Tyr Gly Trp Ser Arg  
1 5 10

<210> 62

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<223> TX125A129E-1

<400> 62

Gln Arg Val Asn Ala Pro Ser Ile Glu Gly Thr Ala Thr  
1 5 10

<210> 63

<211> 14

<212> PRT

<213> Artificial Sequence

<220>

<223> TX-105H-1

<400> 63

Thr Gly Ala Thr Lys His Gly Glu Val Thr Ser Asp Gly Ser  
1 5 10

<210> 64

<211> 12



<212> PRT

<213> Artificial Sequence

<220>

<223> TX-del (123-144)-1r

<400> 64

Gly Ser Ser Arg Arg Gln Thr Arg Tyr Ile Asp Tyr  
1 5 10

<210> 65

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> TX-N1

<400> 65

Gln Thr Ile Gln Pro Gly Thr  
1 5

<210> 66

<211> 22

<212> PRT

<213> Artificial Sequence

<220>

<223> TX-144R-1r

<400> 66

Trp Ala Asn Phe His Asn Ala Thr Asn Val Ser Gly Ser Ser Arg Arg  
1 5 10 15

Asn Arg Arg Val Ser Trp  
20

<210> 67

<211> 15

<212> PRT

<213> Artificial Sequence

<220>

<223> TX-161R-1r

<400> 67

Thr Gly Leu Thr Leu Gly Gln Arg Ala Trp Ala Asn Phe His Asn  
1 5 10 15

<210> 68

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> TX-116G-1

<400> 68

Asp Gly Ser Val Tyr Gly Ile Tyr Arg  
1 5

<210> 69

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

<223> TX-118C-1

<400> 69

Asp Gly Ser Val Tyr Asp Ile Cys Arg Thr Gln Arg  
1 5 10

<210> 70

<211> 13

<212> PRT

<213> Artificial Sequence

<220>

<223> TX-10H11D-1

<400> 70

|     |     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Gly | Thr | Gly | Tyr | His | Asp | Gly | Tyr | Phe | Tyr | Ser | Tyr | Trp |
| 1   |     |     |     | 5   |     |     |     |     | 10  |     |     |     |

<210> 71

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

<223> TX-116G118C-1

<400> 71

|     |     |     |     |     |     |     |     |     |     |     |     |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Asp | Gly | Ser | Val | Tyr | Gly | Ile | Cys | Arg | Thr | Gln | Arg |
| 1   |     |     |     | 5   |     |     |     |     | 10  |     |     |